## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: /0/595, 840
Source: /1540
Date Processed by STIC: 5/24/06

## ENTERED



**IFWO** 

RAW SEQUENCE LISTING DATE: 05/24/2006 PATENT APPLICATION: US/10/595,840 TIME: 15:51:30

Input Set : A:\PTO.KD.txt

Output Set: N:\CRF4\05242006\J595840.raw

```
3 <110> APPLICANT: JEROME, Keith R.
      4 SLOAN, Derek D.
      6 <120> TITLE OF INVENTION: METHODS FOR SCREENING FOR AGENTS CAPABLE OF MODULATING T
LYMPHOCYTE
             FUNCTION IN RESPONSE TO A HERPES SIMPLEX VIRUS-INFECTED CELL
      9 <130> FILE REFERENCE: 14538A-008110US
     11 <140> CURRENT APPLICATION NUMBER: 10/595,840
    12 <141> CURRENT FILING DATE: 2006-05-15
    14 <150> PRIOR APPLICATION NUMBER: PCT/US2004/038050
    15 <151> PRIOR FILING DATE: 2004-11-15
    17 <160> NUMBER OF SEQ ID NOS: 4
    19 <170> SOFTWARE: PatentIn version 3.1
    21 <210> SEQ ID NO: 1
    22 <211> LENGTH: 481
    23 <212> TYPE: PRT
    24 <213> ORGANISM: Herpes simplex virus 1
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    36 His Gly Pro Phe Tyr Thr Pro Ala Glu Asp Ala Cys Leu Asp Ser Pro
                                    40
    40 Pro Pro Glu Thr Pro Lys Pro Ser His Thr Thr Pro Pro Ser Glu Ala
    44 Glu Arg Leu Cys His Leu Gln Glu Ile Leu Ala Gln Met Tyr Gly Asn
                           70
    48 Gln Asp Tyr Pro Ile Glu Asp Asp Pro Ser Ala Asp Ala Ala Asp Asp
    52 Val Asp Glu Asp Ala Pro Asp Asp Val Ala Tyr Pro Glu Glu Tyr Ala
                   100
                                       105
    56 Glu Glu Leu Phe Leu Pro Gly Asp Ala Thr Gly Pro Leu Ile Gly Ala
                                   120
    60 Asn Asp His Ile Pro Pro Pro Cys Gly Ala Ser Pro Pro Gly Ile Arg
    64 Arg Arg Ser Arg Asp Glu Ile Gly Ala Thr Gly Phe Thr Ala Glu Glu
                           150
                                                155
    68 Leu Asp Ala Met Asp Arg Glu Ala Ala Arg Ala Ile Ser Arg Gly Gly
                       165
                                           170
    72 Lys Pro Pro Ser Thr Met Ala Lys Leu Val Thr Gly Met Gly Phe Thr
                   180
                                       185
    76 Ile His Gly Ala Leu Thr Pro Gly Ser Glu Gly Cys Val Phe Asp Ser
               195
                                   200
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80 Ser His Pro Asp Tyr Pro Gln Arg Val Ile Val Lys Ala Gly Trp Tyr

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81		210					215					220				
	ml		mb	0	TT	<b>a</b> 1		7	<b>T</b>			220				
85		ser	Thr	ser		230	Ата	Arg	Leu	Leu	Arg 235	Arg	Leu	Asp	His	Pro 240
		710	T 011	Dwo			7.00	T 011	TT	77-7		C	<b>a</b> 1	**- 7	mla aa	
89	Ala	116	ьец	PIO	245	ьeu	Asp	neu	His	250	vai	ser	GIY		255	Cys
	T 011	17-1	T 011	Dwo		Ф	C1 n	77-	7 ~~		The east	mb	m			7
93	ьeu	vaı	ьеu	260	гуѕ	ıyı	GIII	AIA	Asp 265	Leu	Tyr	Thr	Tyr	ьеи 270	ser	Arg
	7~~	T 011	7 an		LOU	Clu-	71 **~	Dro		т1.	71-	ח ז ה	770 T		7	~1 <u>~</u>
97	Arg	ьeu	275	PIO	ьец	Gry		280	Gln	TTE	Ата	Ald		ser	Arg	GIII
	T 011	T 01	-	- הוג	<b>17-1</b>	7 ~~				7		. al-	285	T1.	. TT-2	Arg
101		290		АІа	. vai	ASL	295 295		: nis	Arg	GII	300		тте	HIS	Arg
				Thr	Glu	Δen			. т.	7.00	The			7.00	т1.	Cys
	305		: шуа	, 1111	Giu	310		FILE	TIE	. ASI	315		, Glu	Asp	116	320
			, Acr	Dho	ദിം			Cvc	Dho	นาไ				7~~		Ser
109		. Gry	Top	FIIC	325		AIG	. Cys	FILE	330		GIY	Ser	ALG		
		Dhe	Dro	Тче			ב ו ג	G1 <sub>3</sub>	r Thr			The	7.00	. או	335	Glu
113		1110	· FIC	340		116	AIG	GLY	345		. ASL	, 1111	ASII	350		Giu
		Τ.Δ11	ב [ג			Dro	Тиг	Thr			. 17-1	7 0 2	. т1-			Ala
117		шси	355		rsp	110	· IYI	360		1111	vai	. nor	365		261	AIA
		Ten			Phe	Glu	ጥ ከተ			Hic	Δer	בומי			Dho	Ser
121	-	370		110	1110	OIU	375		val	1113	noi.	380		пеа	FIIC	Der
				Glv	Pro	Lvs			Pro	Cvs	Δgr			Tle	Thr	Arg
	385		3	017		390		<b>01</b>	110	Cy D	395		0111	. 110		
			Ara	Gln	Δla			His	. Val	Agn			Ser	Pro		Pro
129				0211	405			1110	val	410			. DCI	110	415	
	Glu	Ser	Ara	Leu			Ara	י יי	Ara			r Ala	Δla	Glv		Asn
133			5	420				-1-	425			,		430		11011
	Ara	Pro	Pro			Ara	Pro	Ala			Arc	י דער	ፕν <u>ዮ</u>			Asp
137			435			•		440			2	, -1-	445			
140	Ile	Asp	Val	Glu	Tyr	Leu	Val	Cys	Lvs	Ala	Leu	Thr	Phe	Asp	Glv	Ala
141		450			-		455		•			460			2	
144	Leu	Arq	Pro	Ser	Ala	Ala	Glu	Leu	Leu	Cvs	Leu	Pro	Leu	Phe	Gln	Gln
	465					470				•	475					480
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159	Met	Ala	Cys	Arg	Lys	Phe	Cys	Gly	Val	Tyr	Arg	Arg	Pro	Asp	Lys	Arg
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163	Gln	Glu	Ala	Ser	Val	Pro	Pro	Glu	Thr	Asn	Thr	Ala	Pro	Ala	Phe	Pro
164				20					25					30		
167	Ala	Ser	Thr	Phe	Tyr	Thr	Pro	Ala	Glu	Asp	Ala	Tyr	Leu	Ala	Pro	Gly
168			35		-			40				_	45			-
171	Pro	Pro	Glu	Thr	Ile	His	Pro	Ser	Arg	Pro	Pro	Ser	Pro	Gly	Glu	Ala
172		50					55		_			60				
175	Ala	Arg	Leu	Cys	Gln	Leu	Gln	Glu	Ile	Leu	Ala	Gln	Met	His	Ser	Asp
176						70					75					80

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	<b>~1</b>					•• •	_			<b>a</b> 1.		~-7	~3	~-7	_	~-3
	GIU	Asp	ıyr	Pro		vai	Asp	Ala	Ala		Ala	Glu	Glu	GIu	_	Glu
180	ת 1 ת	λαν	N cm	7 00	85 212	Dwo	7. ~~	7 an	17-1	90 212	П	Dwo	<b>~1</b>	7. ~~	95	21.
184	Ala	ASP	Asp	100	ALA	PIO	Asp	Asp	105	Ala	TYL	PIO	GIU	110	Tyr	Ala
	Glu	Glv	λνα		Lou	Car	Met	1727		λla	Nlα	Dro	T 011		C1	71-
188	Giu	Gry	115	FIIC	пец	SET	Mec	120	PET	ATG	AIA	PIO	125	PIO	Gry	Ala
	Ser	Glv		Dro	Dro	Val.	Pro		λνα	7 T =	בות	Dro		7 cn	v-1	7.20
192	SET	130	nis	FIO	PIO	vai	135	Gry	Arg	Ala	Ата	140	PIO	Asp	vai	AIG
	Thr		Δen	Thr	Glv	Tare	Val	Glv	בומ	Thr	Glv		Thr	Dro	Clu	Glu
	145	Cyb	иор	1111	Gry	150	vai	Gry	AIG	1111	155	FIIG	1111	FIU	GIU	160
		Asp	Thr	Met	Asn		Glu	Δla	T.e11	Ara		Tle	Ser	Δra	Glv	
200	LCu	1101	****		165	9	Olu	niu	шси	170	niu	110	JCI	nr 9	175	Cys
	Lvs	Pro	Pro	Ser		Len	Ala	Lvs	Len		Thr	Glv	Len	Glv		Δla
204	-1-			180				_,_	185			017	<b></b>	190	1110	nia
	Ile	His	Glv		Leu	Ile	Pro	Glv		Glu	Glv	Cvs	Val		Asp	Ser
208			195					200			1	-1	205			
211	Ser	His	Pro	Asn	Tyr	Pro	His		Val	Ile	Val	Lvs		Glv	Trp	Tvr
212		210			•		215					220		2	<b>L</b>	-1-
215	Ala	Ser	Thr	Ser	His	Glu	Ala	Arg	Leu	Leu	Arg	Arg	Leu	Asn	His	Pro
216	225					230		_			235	_				240
219	Ala	Ile	Leu	Pro	Leu	Leu	Asp	Leu	His	Val	Val	Ser	Gly	Val	Thr	Cys
220					245					250					255	
223	Leu	Val	Leu	Pro	Lys	Tyr	His	Cys	Asp	Leu	Tyr	Thr	Tyr	Leu	Ser	Lys
224				260					265					270		
	Arg	Pro		Pro	Leu	Gly	His	Leu	Gln	Ile	Thr	Ala	Val	Ser	Arg	Gln
228			275	_	_			280					285			
	Leu		Ser	Ala	Ile	Asp	Tyr	Val	His	Cys	Lys	_	Ile	Ile	His	Arg
232	_	290	_	1	~3	_	295	_,		_		300		_		_
		iте	ьуs	Thr	GIU		Ile	Pne	тте	Asn		Pro	GIu	Asn	11e	_
	305	<u>ما</u>	7 ~~	Db.	<b>~</b> 1	310	77-	<b>C</b>	D1	**- 7	315		<b>a</b>	3	0	320
240	ьец	GIY	Asp	Pne		Ala	Ala	Cys	Pne		Arg	GIĀ	Cys	Arg		ser
	Dro	Dho	ui a	Фт.	325	т1.	חות	C1	The sec	330	7 ~~	The	7 ~~	77-	335	<b>C1</b>
244	PIO	FIIE	птъ	340	GIY	116	Ala	GIY	345	116	Asp	TIII	ASII	350	PIO	GIU
	Val	T.011	Δla		λen	Dro	Tyr	Thr		17 a 3	Tla	7 cm	T10		Sor	71-
248	·uı	DC u	355	OLY	пор	110	TYL	360	GIII	vai	116	АЗР	365	тър	DCI	AIG
	Glv	Len		Tle	Phe	Glu	Thr		Val	His	Thr	Δla		T.e.11	Phe	Ser
252	0_1	370				014	375		vai			380	001	шси	1110	DCI
	Ala		Ara	Asp	Pro	Glu	Arg	Ara	Pro	Cvs	Asp		Gln	Tle	Ala	Ara
256						390	5	5		-1-	395					400
		Ile	Arq	Gln	Ala		Val	His	Val	Asp		Phe	Pro	Thr	His	
260					405					410					415	
263	Glu	Ser	Arg	Leu	Thr	Ala	His	Tyr	Arq	Ser	Arq	Ala	Ala	Gly		Asn
264			_	420				•	425		_			430		
267	Arg	Pro	Ala	Trp	Thr	Arg	Pro	Ala	Trp	Thr	Arg	Tyr	Tyr	Lys	Ile	His
268			435	_		-		440	_		_	-	445	_		
271	Thr	Asp	Val	Glu	Tyr	Leu	Ile	Cys	Lys	Ala	Leu	Thr	Phe	Asp	Ala	Ala
272		450					455					460				
275	Leu	Arg	Pro	Ser	Ala	Ala	Glu	Leu	Leu	Arg	Leu	Pro	Leu	Phe	His	Pro

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840

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276 465 470 475 480 279 Lys 283 <210> SEQ ID NO: 3 284 <211> LENGTH: 10 285 <212> TYPE: PRT 286 <213> ORGANISM: Homo sapiens 288 <400> SEQUENCE: 3 290 Arg Val Trp Asp Leu Pro Gly Val Leu Lys 294 <210> SEQ ID NO: 4 295 <211> LENGTH: 9 296 <212> TYPE: PRT 297 <213> ORGANISM: Lymphocytic choriomeningitis virus 299 <400> SEQUENCE: 4 301 Lys Ala Val Tyr Asn Phe Ala Thr Cys

VERIFICATION SUMMARY

DATE: 05/24/2006

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